

EXAMINER Monika B. Steinberg

DATE CONSIDERED

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INFORMATION DISCLOSURE CITATION <i>(Use several sheets if necessary)</i>		Docket Number (Optional) 1073.06	Application Number 09/595,096
		Applicant(s) Diller et al.	Filing Date Herewith
*EXAMINER INITIAL <i>MBS</i>	OTHER DOCUMENTS <i>(Including Author, Title, Date, Pertinent Pages, Etc.)</i>		
CC	P.J. Goodford, "A Computational Procedure for Determining Energetically Favorable Binding Sites on Biologically Important Macromolecules", Journal of Medicinal Chemistry, 1985, Vol. 28, No. 7, (1985) American Chemical Society, pp. 849-857.		
CD	C.A. Baxter, C.W. Murray, D.E. Clark, D.R. Westhead & M.D. Eldridge, "Flexible Docking Using Tabu Search and an Empirical Estimate of Binding Affinity", PROTEINS: Structure, Function, and Genetics 33:367-382 (1998), 1998 WILEY-LISS, INC., pp. 367-382.		
CE	G. Jones, P. Willett, R.C. Glen, A.R. Leach & R. Taylor, "Development and Validation of a Genetic Algorithm for Flexible Docking", JMB, J. Mol. Biol. (1997), 267, 727-748, 1997 Academic Press Limited, pp. 726-748.		
CF	M. Rarey, B. Kramer & T. Lengauer, "The Particle Concept: Placing Discrete Water Molecules During Protein-Ligand Docking Predictions", PROTEINS: Structure, Function, and Genetics 34:17-28 (1999), 1999 WILEY-LISS, INC., pp. 17-28.		
CG	B. Kramer, M. Rarey & T. Lengauer, "Evaluation of the FlexX Incremental Construction Algorithm for Protein-Ligand Docking", PROTEINS: Structure, Function, and Genetics 37:228-241 (1999), 1999 WILEY-LISS, INC., pp. 228-241.		
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